

## FIG.2

cleavage site 156-mer 70-mer 5'-226-mer substrate RNA

UG C - GStem II C -G U -A (SEQ ID No. 24)  $U^{GA G-C}$ G<sup>n C</sup> A Stem III Stem I 5'-ACACAACA ACGGGCAC -3' 3'-UGUGUUGUCUGCCCGUG-5' (SEQ ID No. 25) cleavage site

binding with the substrate (secondary prediction map)

B

